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**SEQUENCE LISTING**

<110> AZPIROZ, Ricardo  
CHOE, Sunghwa  
FELDMANN, Kenneth

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Tyr Gly Ser Phe Phe Lys Ser His Ile Leu Gly Cys Pro Thr Ile Val		
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Ser Met Asp Ser Glu Leu Asn Arg Tyr Ile Leu Val Asn Glu Ala Lys		
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Gly Leu Val Pro Gly Tyr Pro Gln Ser Met Ile Asp Ile Leu Gly Lys		

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Pro Tyr Ser Phe Asn Pro Trp Arg Trp Met Asp Lys Ser Leu Glu His		
385	390	395
Gln Asn Ser Phe Leu Val Phe Gly Gly Thr Arg Gln Cys Pro Gly		
405	410	415
Lys Glu Leu Gly Val Ala Glu Ile Ser Thr Phe Leu His Tyr Phe Val		
420	425	430
Thr Lys Tyr Arg Trp Glu Glu Ile Gly Gly Asp Lys Leu Met Lys Phe		
435	440	445
Pro Arg Val Glu Ala Pro Asn Gly Leu Arg Ile Arg Val Ser Ala His		
450	455	460

&lt;210&gt; 21

&lt;211&gt; 444

&lt;212&gt; PRT

&lt;213&gt; Synechocystis sp.

&lt;400&gt; 21

Met Ile Thr Ser Pro Thr Asn Leu Asn Ser Leu Pro Ile Pro Pro Gly

1 5 10 15

Asp Phe Gly Leu Pro Trp Leu Gly Glu Thr Leu Asn Phe Leu Asn Asp

20 25 30

Gly Asp Phe Gly Lys Lys Arg Gln Gln Phe Gly Pro Ile Phe Lys

35	40	45
Thr Arg Leu Phe Gly Lys Asn Val Ile Phe Ile Ser Gly Ala Leu Ala		
50	55	60
Asn Arg Phe Leu Phe Thr Lys Glu Gln Glu Thr Phe Gln Ala Thr Trp		
65	70	75
80		
Pro Leu Ser Thr Arg Ile Leu Leu Gly Pro Asn Ala Leu Ala Thr Gln		
85	90	95
Met Gly Glu Ile His Arg Ser Arg Arg Lys Ile Leu Tyr Gln Ala Phe		
100	105	110
Leu Pro Arg Thr Leu Asp Ser Tyr Leu Pro Lys Met Asp Gly Ile Val		
115	120	125
Gln Gly Tyr Leu Glu Gln Trp Gly Lys Ala Asn Glu Val Ile Trp Tyr		
130	135	140
Pro Gln Leu Arg Arg Met Thr Phe Asp Val Ala Ala Thr Leu Phe Met		
145	150	155
160		
Gly Glu Lys Val Ser Gln Asn Pro Gln Leu Phe Pro Trp Phe Glu Thr		
165	170	175
Tyr Ile Gln Gly Leu Phe Ser Leu Pro Ile Pro Leu Pro Asn Thr Leu		
180	185	190
Phe Gly Lys Ser Gln Arg Ala Arg Ala Leu Leu Leu Ala Glu Leu Glu		
195	200	205
Lys Ile Ile Lys Ala Arg Gln Gln Pro Pro Ser Glu Glu Asp Ala		
210	215	220
Leu Gly Ile Leu Leu Ala Ala Arg Asp Asp Asn Asn Gln Pro Leu Ser		
225	230	235
240		
Leu Pro Glu Leu Lys Asp Gln Ile Leu Leu Leu Leu Phe Ala Gly His		
245	250	255
Glu Thr Leu Thr Ser Ala Leu Ser Ser Phe Cys Leu Leu Leu Gly Gln		
260	265	270
His Ser Asp Ile Arg Glu Arg Val Arg Gln Glu Gln Asn Lys Leu Gln		
275	280	285
Leu Ser Gln Glu Leu Thr Ala Glu Thr Leu Lys Lys Met Pro Tyr Leu		
290	295	300
Asp Gln Val Leu Gln Glu Val Leu Arg Leu Ile Pro Pro Val Gly Gly		
305	310	315
320		
Gly Phe Arg Glu Leu Ile Gln Asp Cys Gln Phe Gln Gly Phe His Phe		
325	330	335
Pro Lys Gly Trp Leu Val Ser Tyr Gln Ile Ser Gln Thr His Ala Asp		
340	345	350
Pro Asp Leu Tyr Pro Asp Pro Glu Lys Phe Asp Pro Glu Arg Phe Thr		
355	360	365
Pro Asp Gly Ser Ala Thr His Asn Pro Pro Phe Ala His Val Pro Phe		
370	375	380
Gly Gly Gly Leu Arg Glu Cys Leu Gly Lys Glu Phe Ala Arg Leu Glu		
385	390	395
400		
Met Lys Leu Phe Ala Thr Arg Leu Ile Gln Gln Phe Asp Trp Thr Leu		
405	410	415
Leu Pro Gly Gln Asn Leu Glu Leu Val Val Thr Pro Ser Pro Arg Pro		
420	425	430
Lys Asp Asn Leu Arg Val Lys Leu His Ser Leu Met		
435	440	

<210> 22  
<211> 519  
<212> PRT  
<213> Zea mays

<400> 22  
 Met Leu Gly Val Gly Met Ala Ala Ala Val Leu Leu Gly Ala Val Ala  
 1 5 10 15  
 Leu Leu Leu Ala Asp Ala Ala Arg Arg Ala His Trp Trp Tyr Arg  
 20 25 30  
 Glu Ala Ala Glu Ala Val Leu Val Gly Ala Val Ala Leu Val Val Val  
 35 40 45  
 Asp Ala Ala Ala Arg Arg Ala His Gly Trp Tyr Arg Glu Ala Ala Leu  
 50 55 60  
 Gly Ala Ala Arg Arg Ala Arg Leu Pro Pro Gly Glu Met Gly Trp Pro  
 65 70 75 80  
 Leu Val Gly Gly Met Trp Ala Phe Leu Arg Ala Phe Lys Ser Gly Lys  
 85 90 95  
 Pro Asp Ala Phe Ile Ala Ser Phe Val Arg Arg Phe Gly Arg Thr Gly  
 100 105 110  
 Val Tyr Arg Ser Phe Met Phe Ser Ser Pro Thr Val Leu Val Thr Thr  
 115 120 125  
 Ala Glu Gly Cys Lys Gln Val Leu Met Asp Asp Asp Ala Phe Val Thr  
 130 135 140  
 Gly Trp Pro Lys Ala Thr Val Ala Leu Val Gly Pro Arg Ser Phe Val  
 145 150 155 160  
 Ala Met Pro Tyr Asp Glu His Arg Arg Ile Arg Lys Leu Thr Ala Ala  
 165 170 175  
 Pro Ile Asn Gly Phe Asp Ala Leu Thr Gly Tyr Leu Pro Phe Ile Asp  
 180 185 190  
 Arg Thr Val Thr Ser Ser Leu Arg Ala Trp Ala Asp His Gly Gly Ser  
 195 200 205  
 Val Glu Phe Leu Thr Glu Leu Arg Arg Met Thr Phe Lys Ile Ile Val  
 210 215 220  
 Gln Ile Phe Leu Gly Gly Ala Asp Gln Ala Thr Thr Arg Ala Leu Glu  
 225 230 235 240  
 Arg Ser Tyr Thr Glu Leu Asn Tyr Gly Met Arg Ala Met Ala Ile Asn  
 245 250 255  
 Leu Pro Gly Phe Ala Tyr Arg Gly Ala Leu Arg Ala Arg Arg Arg Leu  
 260 265 270  
 Val Ala Val Leu Gln Gly Val Leu Asp Glu Arg Arg Ala Ala Arg Ala  
 275 280 285  
 Lys Gly Val Ser Gly Gly Val Asp Met Met Asp Arg Leu Ile Glu  
 290 295 300  
 Ala Gln Asp Glu Arg Gly Arg His Leu Asp Asp Asp Glu Ile Ile Asp  
 305 310 315 320  
 Val Leu Val Met Tyr Leu Asn Ala Gly His Glu Ser Ser Gly His Ile  
 325 330 335  
 Thr Met Trp Ala Thr Val Phe Leu Gln Glu Asn Pro Asp Met Phe Ala  
 340 345 350  
 Arg Ala Lys Ala Glu Gln Glu Ala Ile Met Arg Ser Ile Pro Ser Ser  
 355 360 365  
 Gln Arg Gly Leu Thr Leu Arg Asp Phe Arg Lys Met Glu Tyr Leu Ser  
 370 375 380  
 Gln Val Ile Asp Glu Thr Leu Arg Leu Val Asn Ile Ser Phe Val Ser  
 385 390 395 400  
 Phe Arg Gln Ala Thr Arg Asp Val Phe Val Asn Gly Tyr Leu Ile Pro  
 405 410 415  
 Lys Gly Trp Lys Val Gln Leu Trp Tyr Arg Ser Val His Met Asp Pro  
 420 425 430  
 Gln Val Tyr Pro Asp Pro Thr Lys Phe Asp Pro Ser Arg Trp Glu Gly  
 435 440 445

His Ser Pro Arg Ala Gly Thr Phe Leu Ala Phe Gly Leu Gly Ala Arg  
 450 455 460  
 Leu Cys Pro Gly Asn Asp Leu Ala Lys Leu Glu Ile Ser Val Phe Leu  
 465 470 475 480  
 His His Phe Leu Leu Gly Tyr Lys Leu Ala Arg Thr Asn Pro Arg Cys  
 485 490 495  
 Arg Val Arg Tyr Leu Pro His Pro Arg Pro Val Asp Asn Cys Leu Ala  
 500 505 510  
 Lys Ile Thr Arg Val Gly Ser  
 515

&lt;210&gt; 23

&lt;211&gt; 492

&lt;212&gt; PRT

&lt;213&gt; Danio rerio

&lt;400&gt; 23

Met Gly Leu Tyr Thr Leu Met Val Thr Phe Leu Cys Thr Ile Val Leu  
 1 5 10 15  
 Pro Val Leu Leu Phe Leu Ala Ala Val Lys Leu Trp Glu Met Leu Met  
 20 25 30  
 Ile Arg Arg Val Asp Pro Asn Cys Arg Ser Pro Leu Pro Pro Gly Thr  
 35 40 45  
 Met Gly Leu Pro Phe Ile Gly Glu Thr Leu Gln Leu Ile Leu Gln Arg  
 50 55 60  
 Arg Lys Phe Leu Arg Met Lys Arg Gln Lys Tyr Gly Cys Ile Tyr Lys  
 65 70 75 80  
 Thr His Leu Phe Gly Asn Pro Thr Val Arg Val Met Gly Ala Asp Asn  
 85 90 95  
 Val Arg Gln Ile Leu Leu Gly Glu His Lys Leu Val Ser Val Gln Trp  
 100 105 110  
 Pro Ala Ser Val Arg Thr Ile Leu Gly Ser Asp Thr Leu Ser Asn Val  
 115 120 125  
 His Gly Val Gln His Lys Asn Lys Lys Ala Ile Met Arg Ala Phe  
 130 135 140  
 Ser Arg Asp Ala Leu Glu His Tyr Ile Pro Val Ile Gln Gln Glu Val  
 145 150 155 160  
 Lys Ser Ala Ile Gln Glu Trp Leu Gln Lys Asp Ser Cys Val Leu Val  
 165 170 175  
 Tyr Pro Glu Met Lys Lys Leu Met Phe Arg Ile Ala Met Arg Ile Leu  
 180 185 190  
 Leu Gly Phe Glu Pro Glu Gln Ile Lys Thr Asp Glu Gln Glu Leu Val  
 195 200 205  
 Glu Ala Phe Glu Glu Met Ile Lys Asn Leu Phe Ser Leu Pro Ile Asp  
 210 215 220  
 Val Pro Phe Ser Gly Leu Tyr Arg Gly Leu Arg Ala Arg Asn Phe Ile  
 225 230 235 240  
 His Ser Lys Ile Glu Glu Asn Ile Arg Lys Lys Ile Gln Asp Asp Asp  
 245 250 255  
 Asn Glu Asn Glu Gln Lys Tyr Lys Asp Ala Leu Gln Leu Ile Glu  
 260 265 270  
 Asn Ser Arg Arg Ser Asp Glu Pro Phe Ser Leu Gln Ala Met Lys Glu  
 275 280 285  
 Ala Ala Thr Glu Leu Leu Phe Gly Gly His Glu Thr Thr Ala Ser Thr  
 290 295 300  
 Ala Thr Ser Leu Val Met Phe Leu Gly Leu Asn Thr Glu Val Val Gln  
 305 310 315 320

Lys Val Arg Glu Glu Val Gln Glu Lys Val Glu Met Gly Met Tyr Thr  
           325                 330                 335  
 Pro Gly Lys Gly Leu Ser Met Glu Leu Leu Asp Gln Leu Lys Tyr Thr  
           340                 345                 350  
 Gly Cys Val Ile Lys Glu Thr Leu Arg Ile Asn Pro Pro Val Pro Gly  
           355                 360                 365  
 Gly Phe Arg Val Ala Leu Lys Thr Phe Glu Leu Asn Gly Tyr Gln Ile  
           370                 375                 380  
 Pro Lys Gly Trp Asn Val Ile Tyr Ser Ile Cys Asp Thr His Asp Val  
           385                 390                 395                 400  
 Ala Asp Val Phe Pro Asn Lys Glu Glu Phe Gln Pro Glu Arg Phe Met  
           405                 410                 415  
 Ser Lys Gly Leu Glu Asp Gly Ser Arg Phe Asn Tyr Ile Pro Phe Gly  
           420                 425                 430  
 Gly Gly Ser Arg Met Cys Val Gly Lys Glu Phe Ala Lys Val Leu Leu  
           435                 440                 445  
 Lys Ile Phe Leu Val Glu Leu Thr Gln His Cys Asn Trp Ile Leu Ser  
           450                 455                 460  
 Asn Gly Pro Pro Thr Met Lys Thr Gly Pro Thr Ile Tyr Pro Val Asp  
           465                 470                 475                 480  
 Asn Leu Pro Thr Lys Phe Thr Ser Tyr Val Arg Asn  
           485                 490

&lt;210&gt; 24

&lt;211&gt; 504

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 24

Met Ala Leu Ile Pro Asp Leu Ala Met Glu Thr Trp Leu Leu Leu Ala  
   1                  5                 10                 15  
 Val Ser Leu Val Leu Leu Tyr Leu Tyr Gly Thr His Ser His Gly Leu  
   20                 25                 30  
 Phe Lys Lys Leu Gly Ile Pro Gly Pro Thr Pro Leu Pro Phe Leu Gly  
   35                 40                 45  
 Asn Ile Leu Ser Tyr His Lys Gly Phe Cys Met Phe Asp Met Glu Cys  
   50                 55                 60  
 His Lys Lys Tyr Gly Lys Val Trp Gly Phe Tyr Asp Gly Gln Gln Pro  
   65                 70                 75                 80  
 Val Leu Ala Ile Thr Asp Pro Asp Met Ile Lys Leu Val Leu Val Lys  
   85                 90                 95  
 Glu Cys Tyr Ser Val Phe Thr Asn Arg Glu Pro Phe Gly Pro Val Gly  
   100                105               110  
 Phe Met Lys Ser Ala Ile Ser Ile Ala Glu Asp Glu Glu Trp Lys Arg  
   115                120               125  
 Leu Arg Ser Leu Leu Ser Pro Thr Phe Thr Ser Gly Lys Leu Lys Glu  
   130                135               140  
 Met Val Pro Ile Ile Ala Gln Tyr Gly Asp Val Leu Val Arg Asn Leu  
   145                150               155               160  
 Arg Arg Glu Arg Glu Thr Gly Lys Pro Val Thr Leu Lys Asp Val Phe  
   165                170               175  
 Gly Ala Tyr Ser Met Asp Val Ile Thr Ser Ser Ser Phe Gly Val Asn  
   180                185               190  
 Val Asp Ser Leu Asn Asn Pro Gln Asp Pro Leu Val Glu Asn Thr Lys  
   195                200               205  
 Lys Leu Leu Arg Phe Asp Phe Leu Asp Pro Phe Phe Leu Ser Ile Thr  
   210                215               220

Val Phe Pro Phe Leu Ile Pro Ile Leu Glu Val Leu Asn Ile Cys Val  
 225                   230                   235                   240  
 Phe Pro Arg Glu Val Thr Asn Phe Leu Arg Lys Ala Val Lys Arg Met  
 245                   250                   255  
 Lys Glu Ser Arg Leu Glu Asp Thr Gln Lys His Arg Val Asp Phe Leu  
 260                   265                   270  
 Gln Leu Met Ile Asp Ser His Lys Asn Ser Lys Glu Thr Glu Ser His  
 275                   280                   285  
 Lys Ala Leu Ser Asp Leu Glu Leu Val Ala Gln Ser Ile Ile Phe Ile  
 290                   295                   300  
 Phe Ala Gly Tyr Glu Thr Thr Ser Ser Val Leu Ser Phe Ile Met Tyr  
 305                   310                   315                   320  
 Glu Leu Ala Thr His Pro Asp Val Gln Gln Lys Leu Gln Glu Glu Ile  
 325                   330                   335  
 Asp Ala Val Leu Pro Asn Lys Ala Pro Pro Thr Tyr Asp Thr Val Leu  
 340                   345                   350  
 Gln Met Glu Tyr Leu Asp Met Val Val Asn Glu Thr Leu Arg Leu Phe  
 355                   360                   365  
 Pro Ile Ala Met Arg Leu Glu Arg Val Cys Lys Lys Asp Val Glu Ile  
 370                   375                   380  
 Asn Gly Met Phe Ile Pro Lys Gly Trp Val Val Met Ile Pro Ser Tyr  
 385                   390                   395                   400  
 Ala Leu His Arg Asp Pro Lys Tyr Trp Thr Glu Pro Glu Lys Phe Leu  
 405                   410                   415  
 Pro Glu Arg Phe Ser Lys Lys Asn Lys Asp Asn Ile Asp Pro Tyr Ile  
 420                   425                   430  
 Tyr Thr Pro Phe Gly Ser Gly Pro Arg Asn Cys Ile Gly Met Arg Phe  
 435                   440                   445  
 Ala Leu Met Asn Met Lys Leu Ala Leu Ile Arg Val Leu Gln Asn Phe  
 450                   455                   460  
 Ser Phe Lys Pro Cys Lys Glu Thr Gln Ile Pro Leu Lys Leu Ser Leu  
 465                   470                   475                   480  
 Gly Gly Leu Leu Gln Pro Glu Lys Pro Val Val Leu Lys Val Glu Ser  
 485                   490                   495  
 Arg Asp Gly Thr Val Ser Gly Ala  
 500

<210> 25  
 <211> 575  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Consensus sequence

<221> VARIANT  
 <222> (1)...(575)  
 <223> Xaa = Any Amino Acid or No Amino Acid

<400> 25  
 Xaa  
 1               5               10               15  
 Xaa  
 20              25              30  
 Xaa Xaa Xaa Xaa Xaa Leu Leu Ser Xaa Xaa Ala Leu Xaa Val Xaa  
 35              40              45  
 Leu Xaa Leu Ala Ala Arg Arg Xaa Xaa Xaa Arg Tyr Xaa Xaa Xaa Xaa

50	55	60
Xaa Xaa Xaa Xaa Arg		
Arg Lys Xaa Leu Pro	Pro Gly Thr Met	Gly Leu
65 70	75	80
Pro Xaa Leu Gly Glu	Thr Leu Gln Phe	Leu Lys Xaa Xaa Xaa Xaa
85	90	95
Xaa Pro Gly Asp Phe Xaa Lys	Glu Arg Val Xaa Xaa Tyr	Gly Xaa Xaa
100	105	110
Xaa Xaa Ile Tyr Lys His	Leu Phe Gly Glu Pro	Thr Ile Xaa Ser Xaa
115	120	125
Asp Ala Glu Leu Asn Arg	Phe Xaa Leu Xaa Asn	Glu Gly Xaa Lys Leu
130	135	140
Phe Xaa Cys Xaa Xaa Pro	Ala Ser Xaa Xaa Gly	Xaa Leu Gly Lys Xaa
145	150	155
Ser Leu Xaa Ala Xaa Xaa Gly	Xaa Glu His	Lys Arg Met Arg Xaa Leu
165	170	175
Leu Xaa Ser Xaa Phe	Ser Xaa Xaa Xaa Leu Asp	His Xaa Leu Pro
180	185	190
Xaa Ile Asp Arg Xaa Val	Arg Ser Xaa Leu Xaa Xaa Trp	Xaa Xaa Xaa
195	200	205
Xaa Gln Lys Xaa Xaa Ile	Val Xaa Xaa Xaa Xaa Glu	Xaa Lys Lys Met
210	215	220
Thr Phe Asp Xaa Xaa Xaa Lys	Xaa Xaa Met	Gly Xaa Xaa Pro Xaa Xaa
225	230	235
Glu Xaa Thr Xaa Xaa Xaa Leu	Val Xaa Glu Xaa Glu	Xaa Leu Ile
245	250	255
Lys Gly Leu Phe Ser	Leu Pro Ile Asn	Leu Pro Xaa Thr Ala Tyr Xaa
260	265	270
Lys Ala Leu Xaa Ala Arg	Ala Phe Xaa Xaa Ala	Xaa Leu Glu Xaa Xaa
275	280	285
Xaa Xaa Xaa Xaa Xaa Xaa Xaa Ile	Xaa Glu Xaa Arg	Xaa Glu Glu
290	295	300
Glu Xaa		
305	310	315
Xaa Xaa Xaa Xaa Xaa Asp	Asp Leu	Leu Gly Leu Leu Xaa Ala Xaa
325	330	335
Xaa Xaa Xaa Xaa Glu Asp	Glu Xaa Xaa Xaa Leu	Ser Asp Xaa
340	345	350
Glu Ile Xaa Asp Xaa Ile	Xaa Xaa Leu Leu Phe	Ala Gly His Glu Thr
355	360	365
Thr Ser Ser Xaa Leu Xaa Xaa Ala Val	Lys Phe Leu	Xaa Glu His Pro
370	375	380
Asp Val Xaa Glu Xaa Leu	Arg Glu Glu His	Xaa Ala Ile Xaa Arg Ala
385	390	395
Lys Lys Xaa Xaa Xaa Glu	Ser Xaa Leu Thr	Xaa Xaa Asp Tyr Lys Lys
405	410	415
Met Xaa Tyr Thr Xaa Cys	Val Ile Asn Glu	Thr Leu Arg Leu Ala Xaa
420	425	430
Ile Val Gly Gly Xaa Phe	Arg Xaa Ala Xaa Lys	Asp Val Glu Ile Asn
435	440	445
Gly Tyr Xaa Ile Pro	Lys Gly Trp Lys	Val Xaa Tyr Ser Ile Arg Ala
450	455	460
Val His Leu Asp Pro	Asp Xaa Tyr Pro	Asp Pro Glu Lys Phe Asn Pro
465	470	475
Xaa Arg Trp Xaa Xaa Lys	Xaa Xaa Xaa Ser	Asn Ser Xaa Xaa Xaa
485	490	495
Xaa Xaa Xaa Xaa Xaa Asn	Xaa Xaa Pro	Phe Gly Gly Pro
500	505	510

Arg Leu Cys Pro Gly Lys Glu Leu Ala Lys Leu Glu Met Xaa Val Phe  
515 520 525  
Leu His Arg Leu Val Gln Xaa Phe Trp Glu Leu Ala Xaa Xaa Xaa Asp  
530 535 540  
Xaa Xaa Xaa Lys Leu Val Xaa Phe Pro Thr Xaa Arg Pro Xaa Asp Asn  
545 550 555 560  
Leu Pro Ile Lys Val Xaa Xaa Arg Asp Xaa Xaa Xaa Xaa Xaa Xaa  
565 570 575